(223> Prime (DD)

This could be coused by a bod disk sector.

Per Sequence Rules, (1.824), all submitted

Sequence Listing files must be m' ASUI

Sequence Listing files must be guence Listing

text format. See sample Sequence Listing

(attacked) for valid format.

FITT

<400>

Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

```
Smith, John: Smithgene Inc.
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 <120>
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 <130>
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              PCT/EP98/00001
              1998-12-31
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             1997-10-15
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             1
<210>
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             389
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<212>
<213>
             Paramecium sp.
<220>
<221>
             CDS
             (279)...(389)
<222>
<300>
<301>
             Doe, Richard
<302>
             Isolation and Characterization of a Gene Encoding a
             Protease from Paramecium sp.
<303>
             Journal of Genes
<304>
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<306>
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             1988-06-31
<307>
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                                       ctgggcttct
                                                    caccetgeta
                                                                 atcagatctc
                          cctcttctct
                                                                                E
                                                                                     120
agggagagtg
             tcttgaccct
                          cctctgcctt
                                       tgcagcttca
                                                    caggcaggca
                                                                ggcaggcagc
                                                                                      180
tgatgtggca
                                       cttttcagcc
                                                                 tgggttccgc
             attgctggca
                          gtgccacagg
                                                    aggcttaggg
                                                                                     240
cgcggcgcgg
                                                                cctctcgctc
            cggcccctct
                         cgcgctcctc
                                       togogootot
                                                   ctctcgctct
```

ゴ

Appendix 3, page 2

ggac	ctgat	t aç	ggtga	gcag	gagga	3ggggg	caç	gttago		atg Met l	gtt Val	tca Ser	atg Met	ttc Phe 5	agc Ser	296
ttg Leu	tct Ser	ttc Phe	aaa Lys 10	tgg Trp	cct Pro	gga Gly	ttt Phe	tgt Cys 15	ttg Leu	ttt Phe	gtt Val	tgt Cys	ttg Leu 20	ttc Phe	caa Gln	344
tgt Cys	ccc Pro	aaa Lys 25	gtc Val	ctc Leu	ccc Pro	tgt Cys	cac His 30	tca Ser	tca Ser	ctg -Leu	cag Gln	ccg Pro 35	aat Asn	ctt Leu	: :	389
			٠											i K	<u>:</u> -	
<210 <211		2 37									ì					
<212 <213		PR Pa		ium s	n											
~213	•		20		ρ.											
<400		2		5 1			6	01		91-0	Pro	Clv	Phe	Cua	1.00	
Het 1	Val	Ser	Met	Phe 5	Ser	Leu	Ser	Phe	Lys 10	Trp	PIO	Gly	PHÇ	Cys 15	Leu	
Phe	Val	Cys	Leu 20	Phe	Gln	Cys	Pro	Lys 25	Val	Leu	Pro	Cys	His 30	Ser	Ser	
Leu	Gln	Pro 35	۸sn	Leu		.•										•
<210	•	3				•								•		
<211>		11 PR'	r													
<213>				ial S	equenc	e										
<220> <223>		Des lir	signe nker l	d pepu	ide b	pased o	on si a and	ze and beta	d pola chair	arity ns of	to ac	et as ein XY	a 'Z .			
<400>		3														
Met 1	Val	Λsn	Leu	Glu 5	Pro	Met	His	Thr	Glu 10	Ile						
<210> <400> 000		4														

[Annex VIII follows]

table. The numeric identifier shall be used only in the "Sequence". Listing." The order and presentation of the items of information in the "Sequence Listing" shall conform to the arrangement given below. Each item of information shall begin on a new line and shall begin with the numeric identifier enclosed in angle brackets as shown. The submission of those items of information designated with an "M" is mandatory. The submission of those items of information designated with an "O" is optional. Numeric identifiers <110> through <170> shall only be set forth at the beginning of the "Sequence Listing." The following table illustrates the numeric identifiers.

Numeric Identifier	Definition	Comments and Format	Mandatory (M) or Optional (O)
<110>	Applicant	Preferably max. of 10 names; one name per line; preferable format: Surname, Other Names and/or Initials	M · · ·
<120>	Title of Invention		М
<130>	File Reference	Personal file reference	M. when filed prior to assignment of appl. number
<140>	Current Applica- tion Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if available
<141>	Current Filing Date	Specify as: yyyy-mm-dd	M, if available
<150>	Prior Application Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if applicable include priority documents under 35 USC 119 and 120
<151>	Prior Application Filing Date	Specify as: yyyy-mm-dd	M, if applicable
<160>	Number of SEQ ID NOs	Count includes total number of SEQ ID NOs	М
<170>	Softwäre	Name of software used to create the . Sequence Listing	0
<210>	SEQ ID NO:#:	Response shall be an integer representing the SEQ ID NO shown	M
<211>	Length	Respond with an integer expressing the number of bases or amino acid residues	M

<212>

Type

Whether presented sequence molecule is DNA, RNA, or PRT (protein). If a nucleotide sequence contains both DNA and RNA fragments, the type shall be "DNA." In addition, the combined DNA/ RNA molecule shall be further

described in the <220> to <223> feature

section.

<213> Organism

Scientific name, i.e. Genus/species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Sequence" organisms shall be further described in the <220> to <223>

<220> Feature

Leave blank after <220>. <221-223> provide for a description of points of biological significance in the sequence.

feature section.

M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGAN-ISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA.

М

<221>

Name/Key

Provide appropriate identifier for feature, preferably from WIPO Standard ST.25 (1998), Appendix 2, Tables 5 and 6

M, under the following conditions:⊨ if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence

<222> Location Specify location within sequence; where appropriate state number of first and last bases/amino acids M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified

•			·			
•		in feature	base was used in a sequence			
<223>	Other Infor- mation	Other relevant information; four lines maximum	M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if			
			ORGANISM is "Artificial Sequence" or "Unknown"; if molecule is com- bined DNA/RNA.			
<300>	Publication Information	Leave blank after <300>	0			
<301>	Authors	Preferably max of ten named authors of publi- cation; specify one name per line; preferable format: Surname, Other Names and/or Initials	0			
.2025	mi n l -	INICIAIS	0			
<302>	Title					
<303>	Journal		0			
<304>	Volume		0			
<305>	Issue		0			
<306>	Pages		0			
<307>	Date	Journal date on which data published; specify as yyyy-mm-dd, MMM-yyyy or Season-yyyy				
<308>	Database Accession Number	Accession number assigned by data-base including database name	0			
<309>	Database Entry Datė	Date of entry in database; specify as yyyy-mm-dd or MMM-yyyy	0			
<310>	Patent Document Number	Document number; for patent-type citations only. Specify as, for example, US 07/999,999	0			

<311>	Patent Filing Date	Document filing date, for patent-type citations only; specify as yyyy-mm-dd	0
<312>	Publication Date	Document publication date, for patent-type citations only; specify as yyyy-mm-dd	0
<313>	Relevant Residues	FROM (position) TO (position)	0
<400>	Sequence	SEQ ID NO should follow the numeric identifier and should appear on the line preceding the actual sequence	м .

- 5. Section 1.824 is revised to read as follows:
- 1.824 Form and format for nucleotide and/or amino acid sequence submissions in computer readable form.
- (a) The computer readable form required by 1.821(e) shall meet the following specifications:
- (1) The computer readable form shall contain a single "Sequence Listing" as either a diskette, series of diskettes, or other permissible media outlined in paragraph (c) of this section.
- (2) The "Sequence Listing" in paragraph (a) (1) of this section shall be submitted in American Standard Code for Information Interchange (ASCII) text. No other formats shall be allowed.
- (3) The computer readable form may be created by any means, such as word processors, nucleotide/amino acid sequence editors or other custom computer programs; however, it shall conform to all specifications detailed in this section.
- (4) File compression is acceptable when using diskette media, so long as the compressed file is in a self-extracting format that will decompress on one of the systems described in paragraph (b) of this section.
- (5) Page numbering shall not appear within the computer readable form version of the "Sequence Listing" file.
- (6) All computer readable forms shall have a label permanently affixed thereto on which has been hand-printed or typed: the name of the applicant, the title of the invention, the date on which the data were recorded on the computer readable form, the operating system used, a reference number, and an application serial number and filing date, if known.
- (b) Computer readable form submissions must meet these format requirements:
- (1) Computer: IBM PC/XT/AT, or compatibles, or Apple Macintosh;
- (2) Operating System: MS-DOS, Unix or Macintosh;